

Dementias in the Genomic Era:

Current Understanding of Frontotemporal Dementia

Demystifying Medicine Lecture
NIH Campus, Bethesda, MD
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A large, semi-transparent circular graphic on the left side of the slide features a stylized brain circuit board logo. To its right is a block of multi-colored DNA sequence text. The text is composed of various nucleotide bases (A, T, C, G) in different colors (red, green, blue, yellow), representing a genomic sequence. The sequence is extremely long and dense, spanning the width of the slide area.

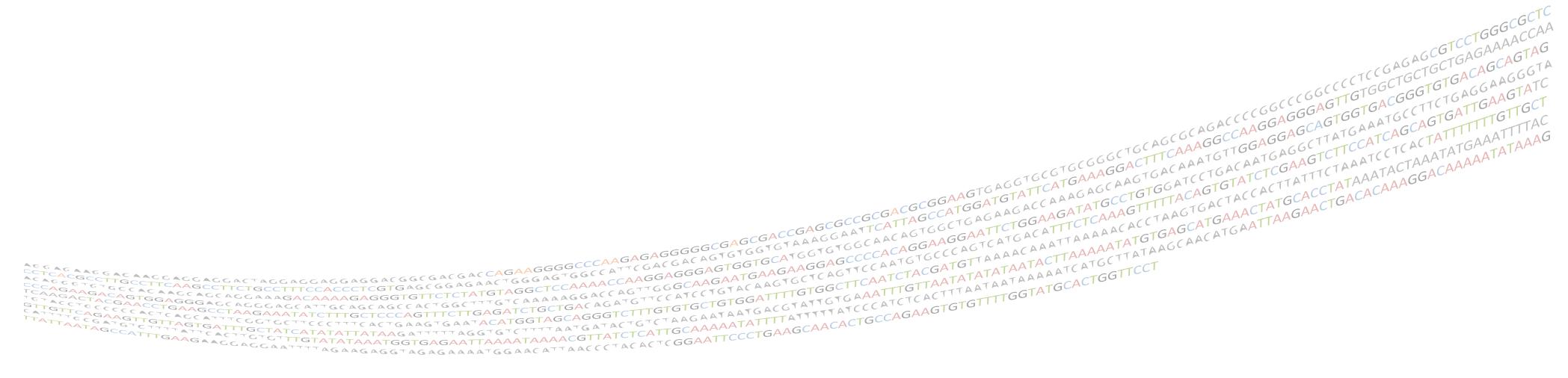


National Institutes of Health
Turning Discovery Into Health

Disclosures

- Patents on the clinical testing and therapeutic interventions for the hexanucleotide repeat expansion of *C9orf72*

- Funding from Merck, CDC, Myasthenia Gravis Foundation, ALSA, Packard Center



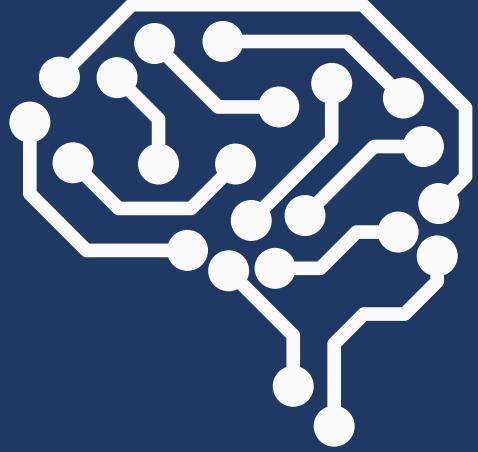
A decorative graphic at the bottom of the slide features several overlapping DNA double helix molecules. The DNA strands are composed of various colored nucleotides (A, T, C, G) and are oriented diagonally across the page. The colors of the nucleotides include green, blue, red, yellow, and purple, creating a complex, overlapping pattern that spans from the bottom left to the top right of the slide area.

Learning Objectives

- To understand the clinicopathological features of frontotemporal dementia
- To review current genetic understanding of frontotemporal dementia
- To delineate how genetic knowledge can be leveraged to advance the precision medicine paradigm

The watermark consists of a dense, multi-colored sequence of DNA base pairs (A, T, C, G) arranged in a grid pattern, representing a genome sequence.

Overview



1

Background

2

Illustrative Cases

3

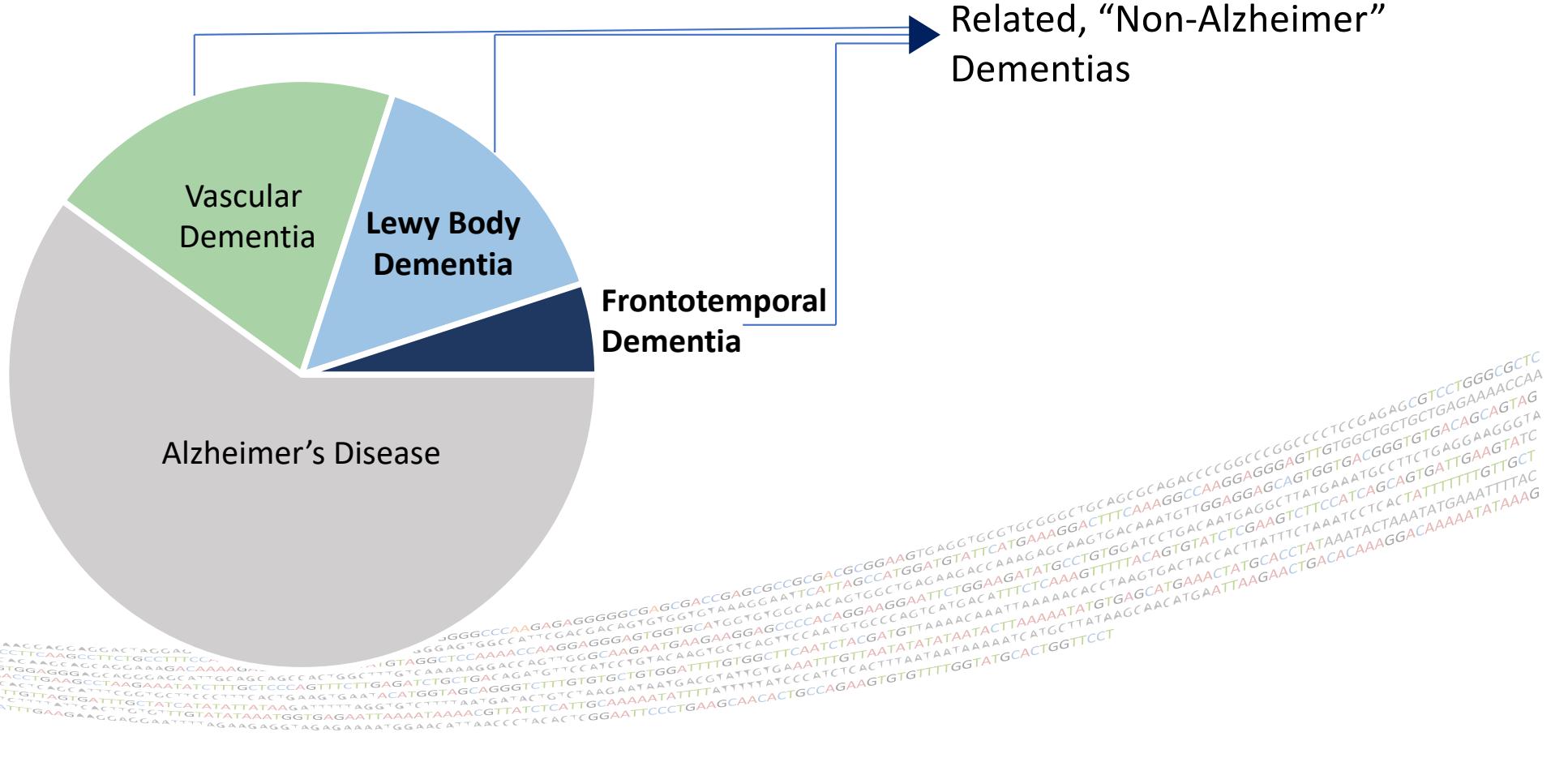
Current Genetics

4

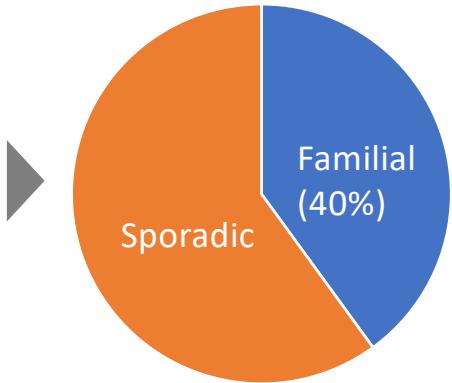
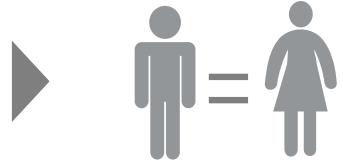
Future Directions



Age-Related Dementias



Epidemiology of frontotemporal dementia



- Prevalence 15-20 per 100,000 population
- 20,000 – 30,000 cases in the U.S.
- Incidence 3.5 per 100,000 population per year
(age range 45-64 years)

Onyike, CU et al. *Int Rev Psychiatr*, (2013)
Rohrer, JD et al. *Neurology* (2009)

A Historic View on FTD

1892



1851 - 1924

71 year-old man with

- progressive cognitive decline
- personality change
- speech disorder: *impaired comprehension*
paraphasias
partial preservation of repetition

Asymmetric temporal atrophy on pathology

Pick A., Ueber die Beziehungen der senilen Hirnatrophie zur Aphasie, *Prag Med Wochenschr*, 17, 165-7 (1892)
Image source: www.wikipedia.org

A Historic View on FTD

1892 1911

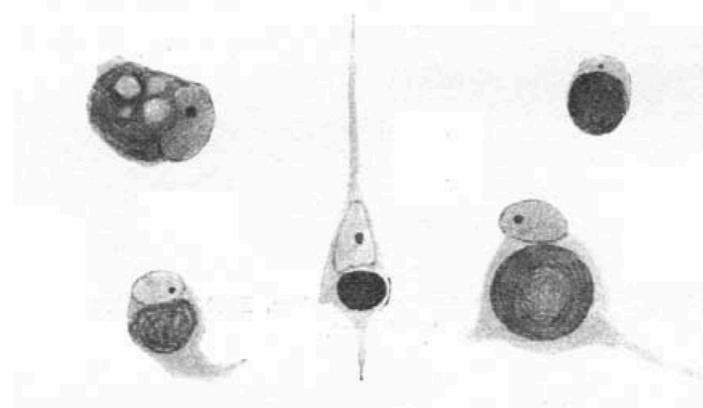


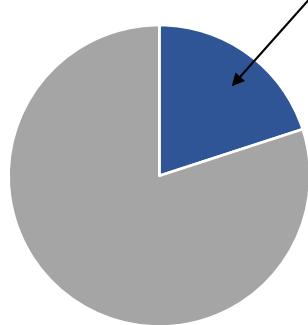
Illustration of argyophilic Pick bodies by Alois Alzheimer. (*Z Gesamte Neurol Psychiatr* 1911; 4: 356–85).

Image source:
www.wikipedia.org

A Historic View on FTD

Timeline: 1892, 1956, 1974

Only 20% of cases have Pick bodies



First pathological classification

Escourrolle, R. *La maladie de Pick. Etude D'ensemble et Synthese Anatomico-Clinique*, (1956)
Constantinidis, J. et al. *Eur Neurol* (1974)

A Historic View on FTD



Genetic studies reveal disease-causing mutations

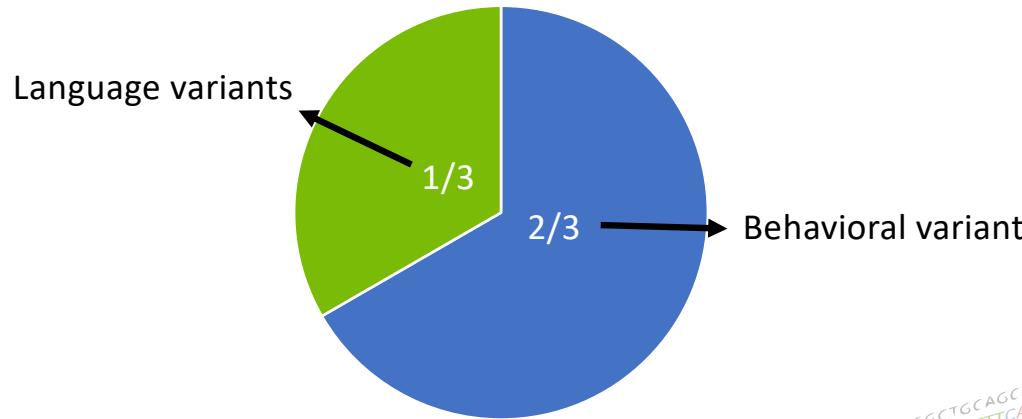
Immunohistochemistry: diverse protein aggregations

Escourroule, R. *La maladie de Pick. Etude D'ensemble et Synthese Anatomico-Clinique*, (1956)
Constantinidis, J. et al. *Eur Neurol* (1974)

A dense, colorful sequence of DNA base pairs (A, T, C, G) arranged in a spiral pattern, representing the genetic code. The colors used are blue, red, green, and yellow, corresponding to the four nucleotide bases.

Clinical FTD Subtypes

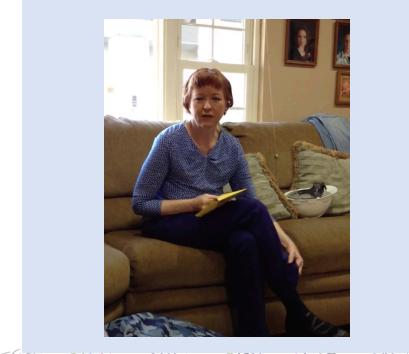
FRONTOTEMPORAL DEMENTIA



Clinical Features of FTD

FRONTOTEMPORAL DEMENTIA

Behavioral variant	Language variant (PPA)	Non-fluent variant	Semantic variant
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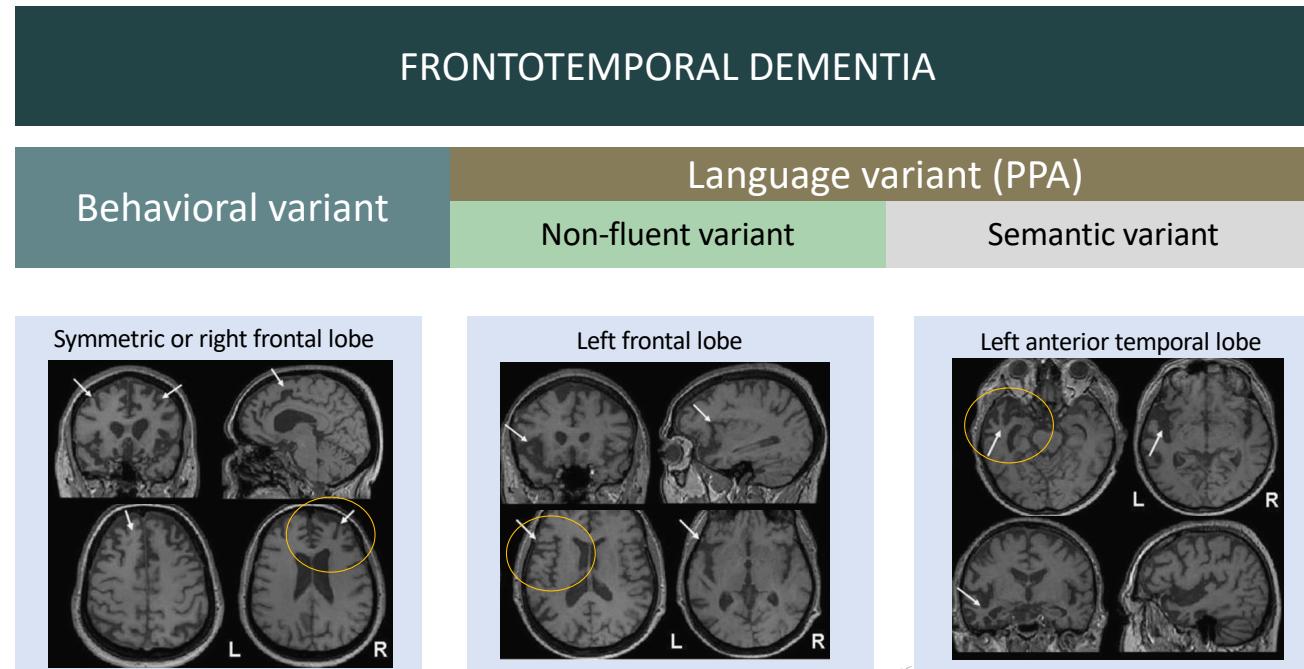


Source: <https://www.youtube.com/watch?v=14bWxelvshc>

Source: <https://www.youtube.com/watch?v=eYUGzChNmNA>

The slide features a grid of four sections. The top section is a dark green box containing the text "FRONTOTEMPORAL DEMENTIA". Below it is a 2x2 grid of colored boxes: a teal box on the left labeled "Behavioral variant", a gold box at the top labeled "Language variant (PPA)", a light green box in the middle labeled "Non-fluent variant", and a light grey box on the right labeled "Semantic variant". Below the grid are two photographs: one of a man in a red cap and another of a woman on a couch. To the right of the photographs is a large, tilted block of multi-colored text, likely representing a speech or video transcription. The text is in various colors (red, blue, green, yellow) and is rotated diagonally for readability.

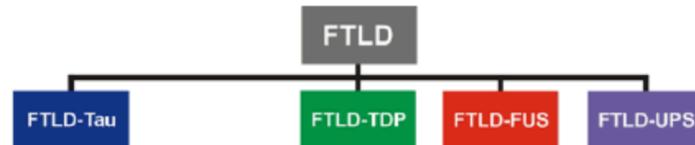
Neuroimaging in Clinical FTD Subtypes



Acosta, F. et al. Cortex (2012)

Abnormal Protein Aggregation

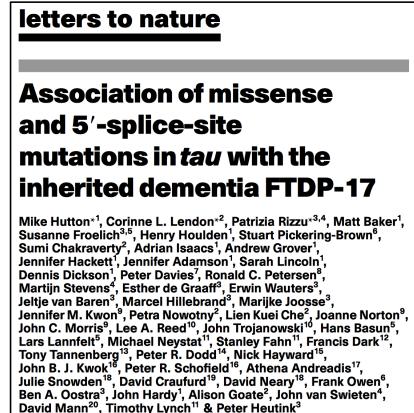
FRONTOTEMPORAL LOBAR DEGENERATION



Jellinger K, *Journal of Alzheimer disease and Parkinsonism*, 2017

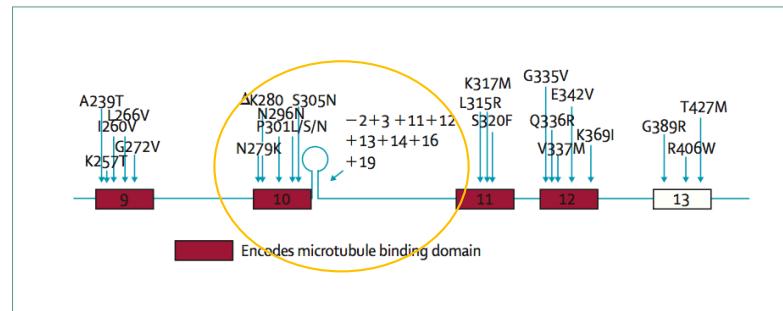
Genetics of FTD: *MAPT*

1892



1998

10% of familial FTD cases



Missense & splice-site mutations

Gain of function

Neary, D. et al. *Lancet Neurol* (2005)

Tauopathies

Progressive Supranuclear Palsy

Corticobasal Degeneration

Alzheimer Dementia

Frontotemporal Dementia

Nodding Syndrome

Parkinson-Dementia Complex of Guam

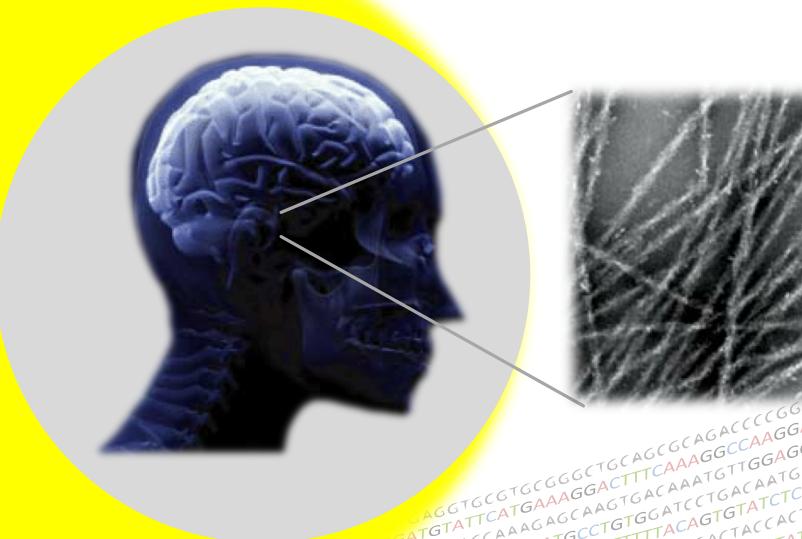


Image source: Max Planck Society

The Genetics of FTD: *GRN*

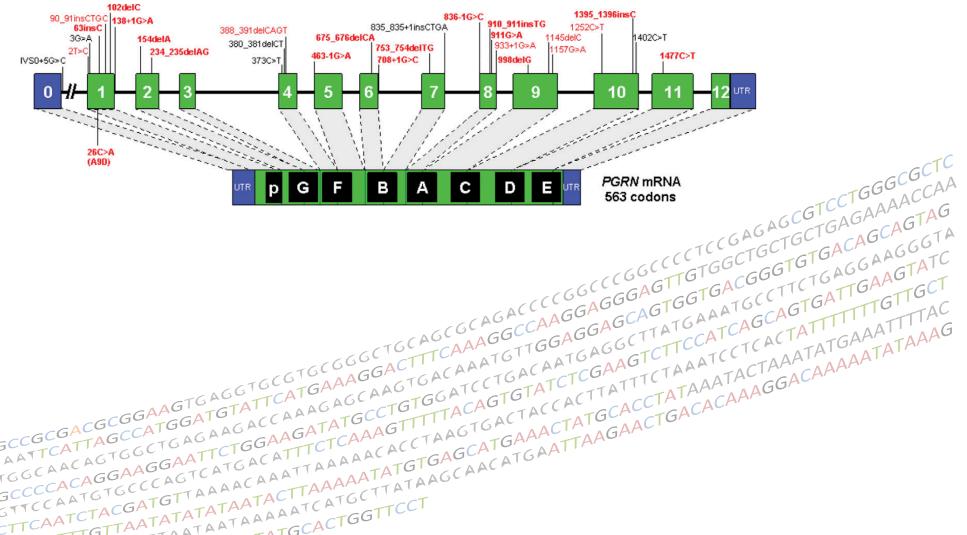
1892

2006

10% of familial cases

Heterozygous missense mutations

Haploinsufficiency



Gass, J et al., Hum Mol Genet (2006)

The Genetics of FTD: *C9orf72*

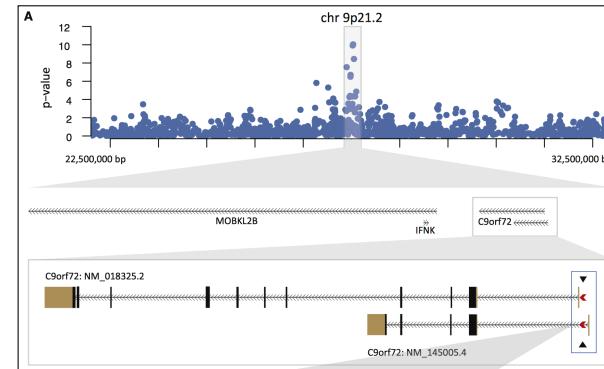
1892

2011

30% of familial cases

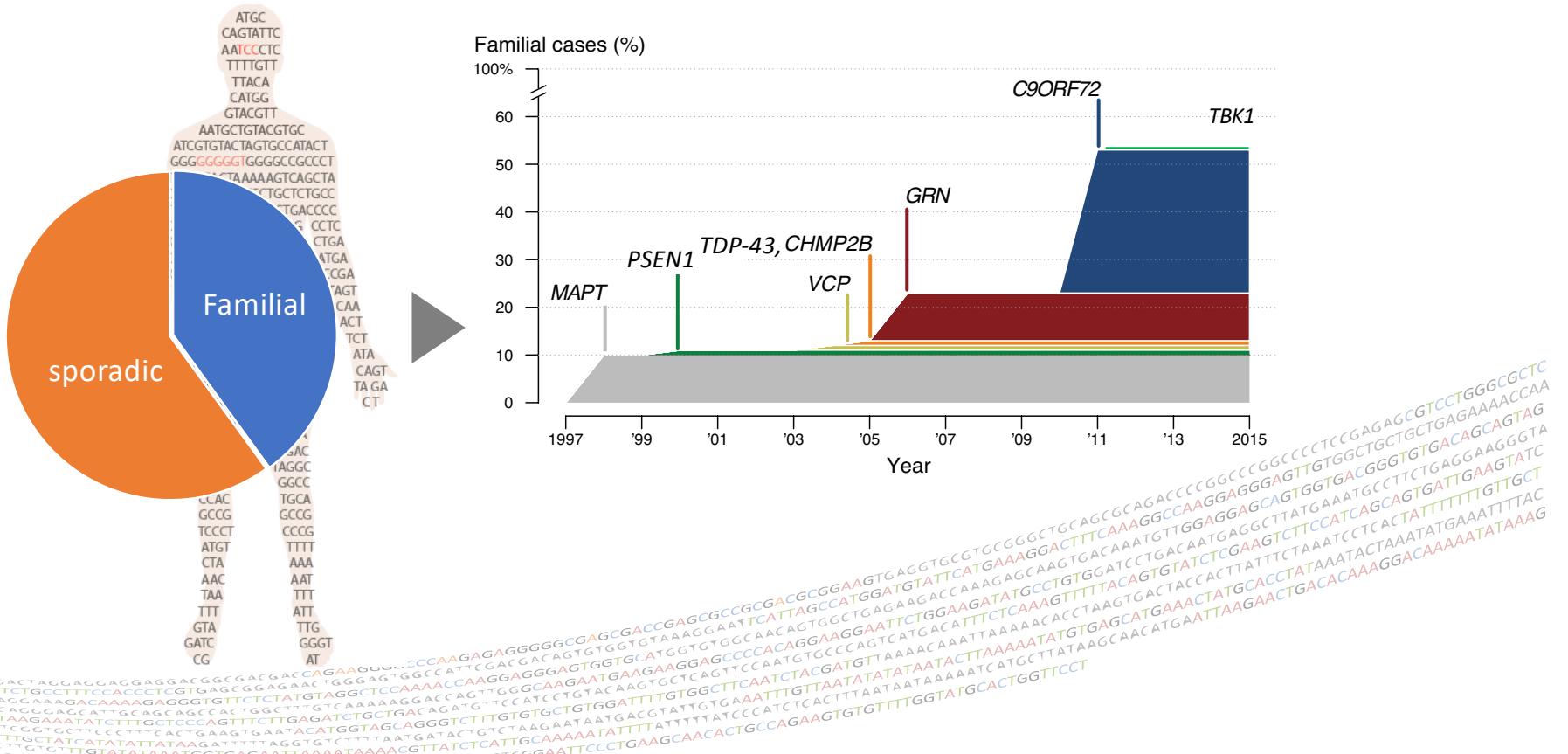
Hexanucleotide repeat expansion
[GGGGCC] n x 700-1600 (nr: <30 repeats)

Molecular link between FTD & ALS

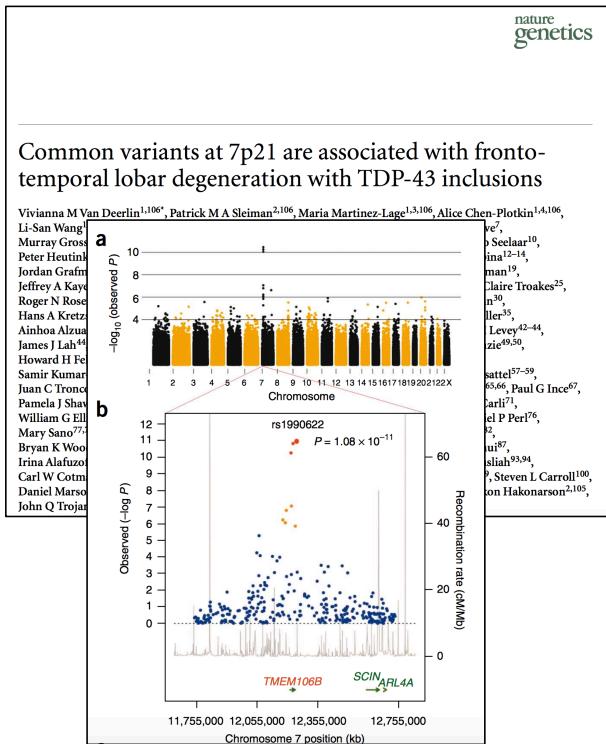


Renton, A et al., *Neuron* (2011)
DeJesus-Hernandez, et al., *Neuron* (2011)

The Genetics of FTD



FTD GWAS: a Role for Common Variants?



N=515 FTD-TDP patients, 2509 controls

Replication: 89 cases, 553 controls

European ancestry FTD-TDP cases

Increased expression of TMEM106B

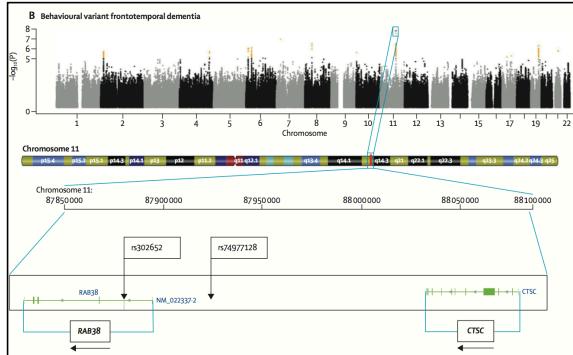
Lysosomal metabolism

Van Deerlin, V et al. *Nature Genetics* (2010)

FTD GWAS: a Role for Common Variants?

Frontotemporal dementia and its subtypes: a genome-wide association study

Raffaele Ferrari*, Dena G Hernandez*, Michael A Nalls*, Jonathan D Rohrer*, Adaikalanay Ramasamy, John BJ Kwok, Carol Dobson-Stone, William S Brooks, Peter R Schofield, Clenda M Halliday, John R Hodges, Oliver Piguet, Lauren Bartley, Elizabeth Thompson, Eric Haan, Isabel Hernández, Agustín Ruiz, Merce Boada, Barbara Borroni, Alessandro Padovani, Carlos Cruceanu, Nigel J Cairns, Luisa Benussi, Giuliano Binetti, Roberta Ghidoni, Gianluigi Forloni, Daniela Galimberti, Chiara Feruglio, Maria Serpetzis, Elio Scarpini, Jordi Clarimón, Alberto Lledó, Rafael Bleas, María Landínez Walldé, Karin Nilsson, Christer Nilsson, Jan R Mackenzie, Ging-Yuek R Hsiung, David M A Mann, Jordan Grafman, Christopher M Morris, Johannes Attems, Timothy D Griffiths, Jan G McKeith, Alan J Thomas, P Pietrini, Edward D Huyck, Eric M Wassermann, Attila Baboci, Evelyn Jaros, Michael C Tierney, Paul Pastor, Cristina Razquin, Sara Ortego-Cubero, Elena Alonso, Robert Perneczky, Jönne Diehl-Schmid, Panagiotis Alexopoulos, Alexander Kurz, Innocenzo Rainiero, Elisa Rubino, Lorenzo Pinelli, Ekaterina Rogava, Peter S George-Hylop, Giacomo Rossi, Fabrizio Taglioni, Giorgia Giacomini, James B Rose, Johannes C M Schlaichetzki, James Uphill, John Collinge, Simon Mead, Adrian Danek, Vivianne M Van Derlin, Murray Grossman, John Q Trojanowski, Julie van der Zee, William Deschamps, Tim Van Langenhove, Marc Cruts, Christine Van Broeckhoven, Stefano F Cappa, Isabelle le Ber, Didier Hennequin, Véronique Gilje, Martine Verelletta, Aleix Brice, Benedetta Nascim, Sandra Sorbi, Silvia Baglioni, Irene Placeri, Jürgen E Nielsen, Lena E Herník, Matthias Riemschneider, Manuel Mayhous, René Debat, Gilles Gasparoni, Sabrina Pichler, Wei Gu, Martin N Rossor, Nick C Fox, Jason D Warren, Maria Grazia Spillantini, How R Morris, Patrizia Lucucci, Peter Hockin, Julie S Snedden, Sara Bellonson, Anna Richardson, Alexandre Gerhard, Anna C Brunni, Raffaele Maletta, Francesca Frangione, Chiara Cupidi, Livio Bernardi, Maria Anfossi, Mauro Gallo, Maria Elena Conidi, Nicoletta Smeraldi, Rosalba Badanesco, Matt Baker, Dennis W Dickson, Neill R Goffe-Rodriguez, Ronald C Petersen, David Knopman, Keith A Josephs, Bradley F Boeve, Joseph F Parisi, William W Seeley, Bruce L Miller, Anna M Karydas, Howard Rosen, John C van Swieten, Elise G Doppler, Hanno Selvaggi, Yolande A L Pijnenburg, Philip Scheltens, Giancarlo Logroscino, Rosa Capozzo, Valeria Novelli, Annibale A Puccia, Massimo Fratiglione, Alfredo Postiglione, Graziella Milani, Paolo Sorrentino, Mark Kristiansen, Huei-Hsin Chang, Caroline Graff, Florence Pasquier, Adeline Rollin, Vincent Deramecourt, Florence Lebert, Dimitrios Kapogiannis, Luigi Ferrucci, Stuart Pickering-Brown, Andrew B Singleton, John Hardy, Parastoo Momeni



N=2154 FTD patients, 4308 controls

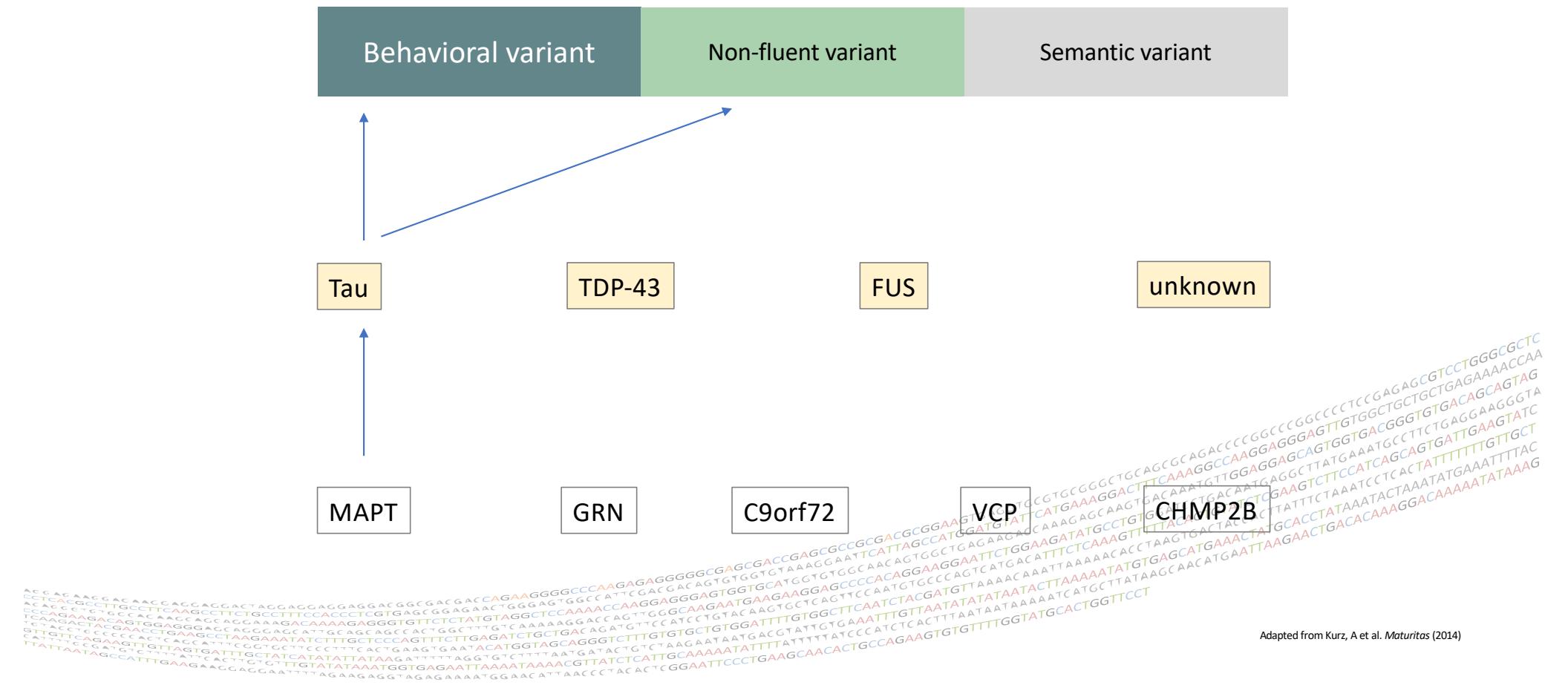
European ancestry FTD cases

RAB38/CTSc in bvFTD

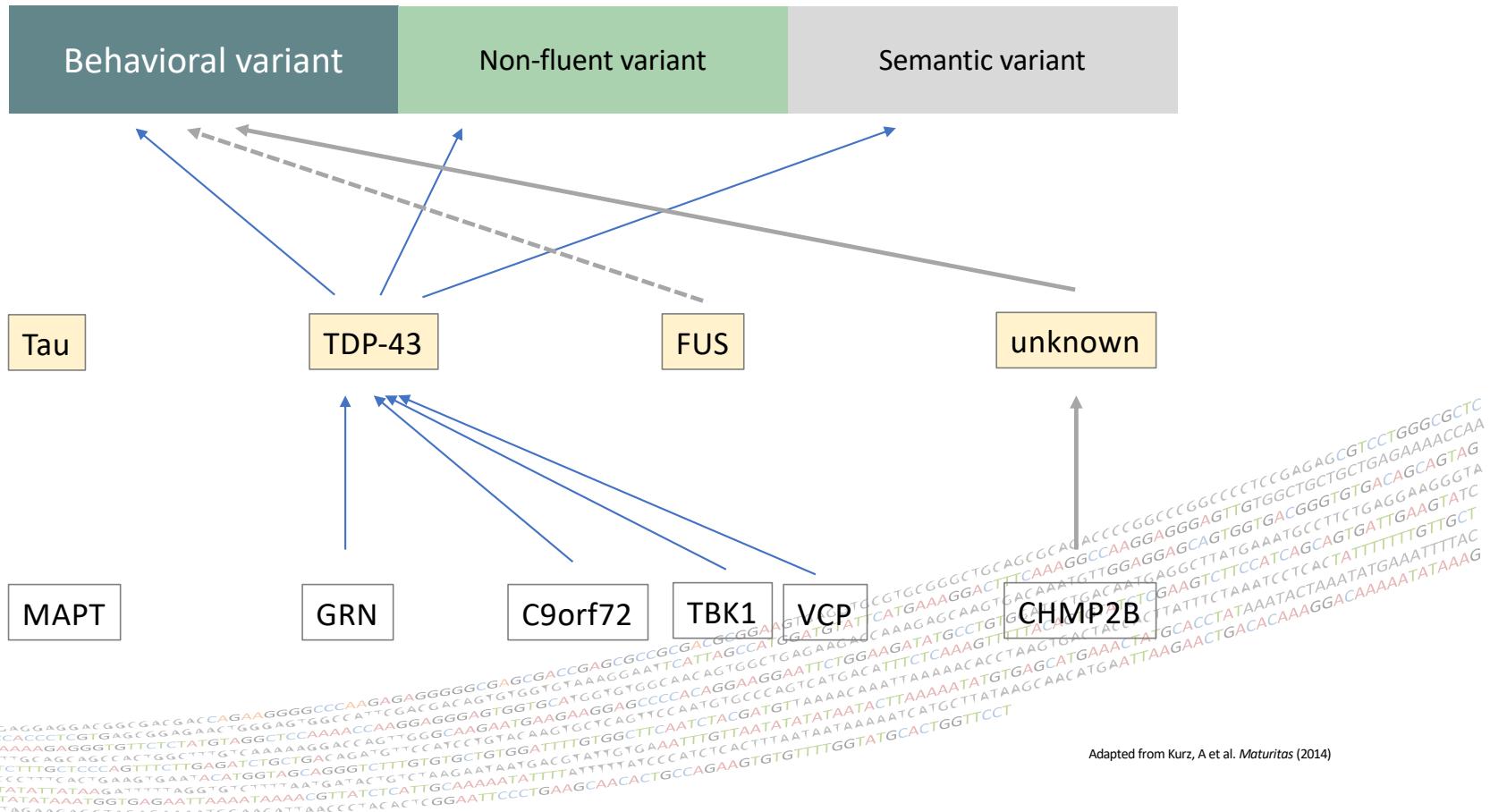
Lysosomal metabolism

Ferrari, R et al. *Lancet Neurology* (2014)

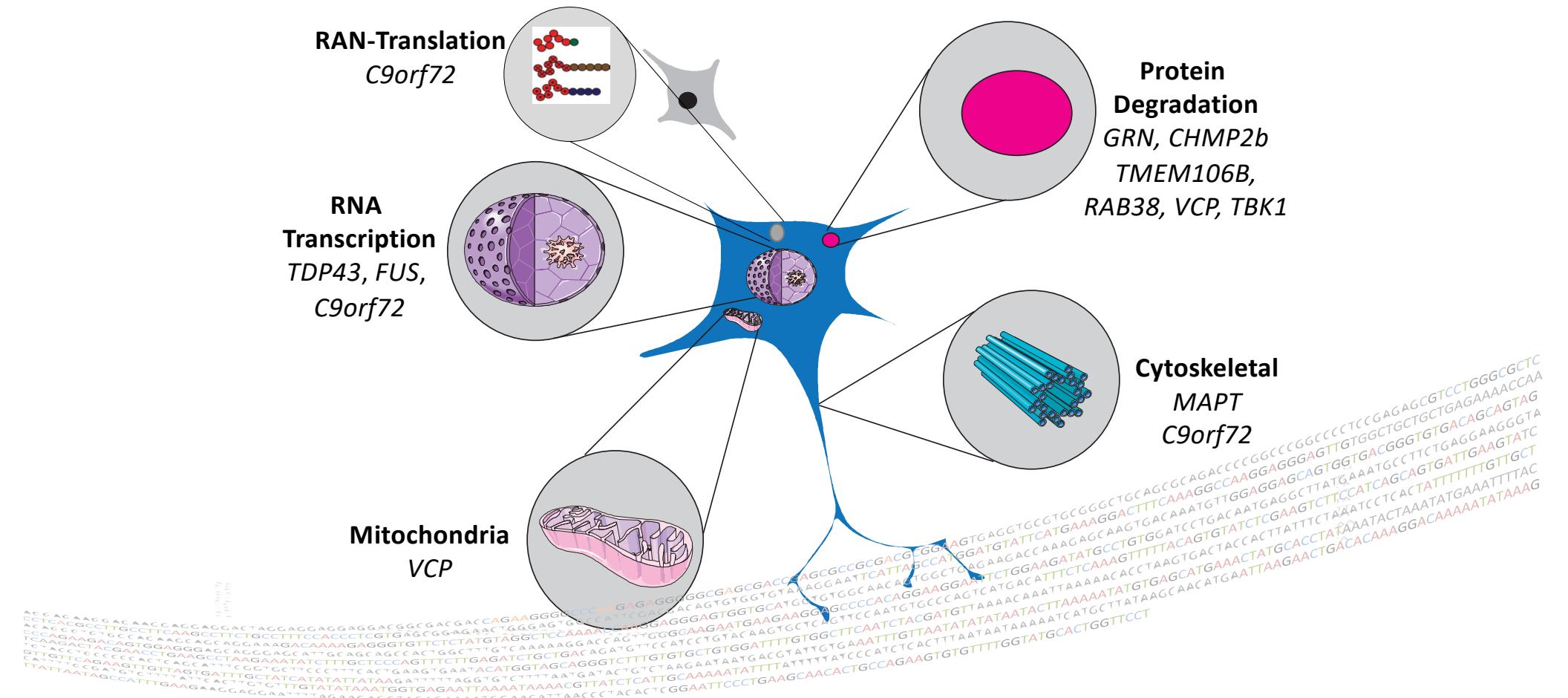
Genotype-Phenotype Correlations



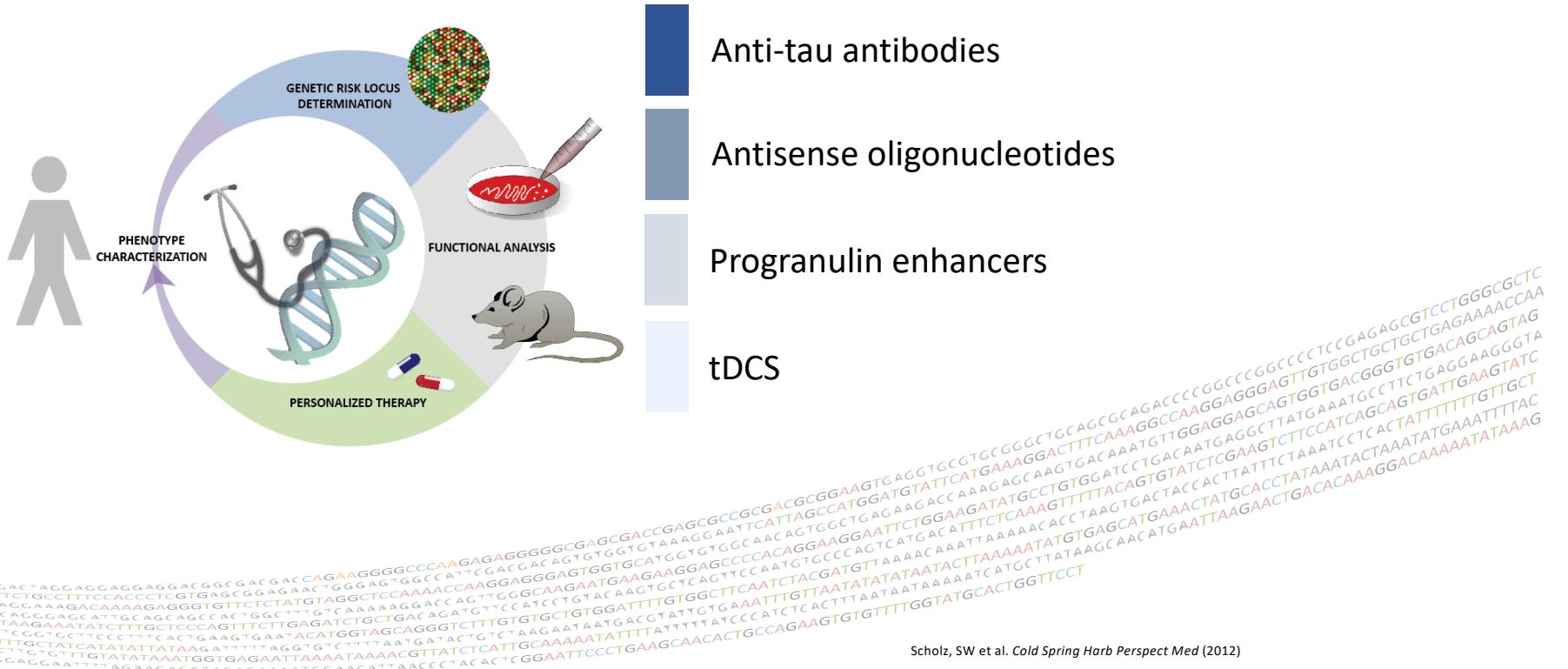
Genotype-Phenotype Correlations



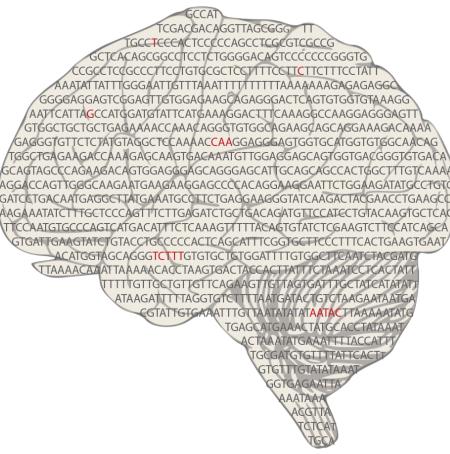
Pathogenic Mechanisms of FTD



Future Directions in FTD: Disease Modification



Summary



Common form of dementia

Clinically, pathologically and genetically heterogeneous

Treatment is supportive

Pathogenesis is poorly understood

